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mask repetitive contig sequence unique

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Analysis of major **repetitive** DNA sequences in the dog(*Canis familiaris*) genome

S Bentolila, JM Bach, JL Kessler, I Bordelais & ... - Mammalian Genome, 1999 - springerlink.com

... the analysis of DNA sequences (repeat **mask**- ing, definition of ... 270 Huang X (1992)A **contig** assembly program ... Stiegler GL (1992) A highly **repetitive** DNA **sequence** ...Cited by 17 - Web Search - ncbi.nlm.nih.gov - ncbi.nlm.nih.govSearching the Expressed **Sequence** Tag (EST) Databases: Panning for Genes

CV Jongeneel - Briefings in Bioinformatics, 2000 - ingentaconnect.com

... from several species, under the denomination of **unique** gene indices. ... to return gapped alignments between the query and the database **sequence**. ... **Contig** databases ...Cited by 12 - Web Search - ismb00.sdsc.edu - sib-dea.unil.ch - amber.cs.umd.edu - all 9 versions »Gene discovery by EST sequencing in *Toxoplasma gondii* reveals sequences restricted to the

JW Ajioka, JC Boothroyd, BP Brunk, A Hehl, L ... - Genome Res, 1998 - genome.org

... As in (3), above, **repetitive** elements were identified ... Lipman, unpubl.) was used to**mask** low entropy ... **sequence** was determined for each **contig** defined (available ...Cited by 75 - Web Search - genome.org - path.cam.ac.uk - ncbi.nlm.nih.gov - all 7 versions »Comparative Genome Mapping in the **Sequence**-based Era: Early Experience with Human Chromosome 7

JW Thomas, TJ Summers, SQ Lee-Lin, VV Maduro, JR ... - Genome Research, 2000 - genome.org

... of a mouse **sequence**-ready BAC **contig** using human ... Roughly 180 kb of genomic **sequence** (top line ... UWGC/analysistools/repeatmask.htm) to **mask** **repetitive** elements and ...Cited by 21 - Web Search - csb.yale.edu - pubmedcentral.nih.gov - dx.doi.org - all 8 versions »Genomic organization of the 260 kb surrounding the waxy locus in a Japonica rice

H Nagano, L Wu, S Kawasaki, Y Kishima, Y Sano - Genome, 1999 - article.pubs.nrc-cnrc.gc.ca

... repeti- tive short **sequence**(s), which can **mask** their own ... the single-copy regions in the 260-kb **contig** are interrupted by **repetitive** sequences every ...Cited by 7 - Web Search - dx.doi.org - pubs.nrc-cnrc.gc.ca - ncbi.nlm.nih.gov - all 6 versions »The Mouse Clock Locus: **Sequence** and Comparative Analysis of 204 Kb from Mouse Chromosome 5

LD Wilsbacher, AM Sangoram, MP Antoch, JS ... - Genome Research, 2000 - genome.org

... 11-22 align within the largest **contig**, while exons ... of introns as well (mouse **repetitive** elements were ... is possible that position effects could **mask** a regulatory ...Cited by 4 - Web Search - pubmedcentral.nih.gov - dx.doi.org - ncbi.nlm.nih.gov - all 6 versions »Identification of a gene that reverses the immortal phenotype of a subset of cells and is a member ...

MJ Bertram, NG Berube, X Hang-Swanson, Q Ran, JK ... - Mol. Cell. Biol, 1999 - mcb.asm.org

... cDNAs containing **repetitive** elements were eliminated from further ... have coding **sequence** within the **contig** (1). One ... cell in the population to **mask** the senescence ...Cited by 60 - Web Search - mcb.asm.org - pubmedcentral.nih.gov - ncbi.nlm.nih.gov - all 5 versions »Data mining parasite genomes: haystack searching with a computer

D LAWSON - Parasitology, 1999 - journals.cambridge.org

... The generation of assembled **contig** consensus from the primary ... regions of low- complexity can **mask** true hits ... genome or the degree of **repetitive** sequences present ...
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Two regions of homozygous deletion clusters at chromosome band 9 p 21 in human lung cancer

K Hamada, T Kohno, M Takahashi, M Yamazaki, H ... - Genes Chromosomes and Cancer, 2000 - [doi.wiley.com](#)
 ... After **mask**- ing of these **repetitive** sequences, potential exons ... estimated as being more than 2 Mb in a YAC **contig** map (Povey et al., 1994), and the distance ...
 Cited by 14 - [Web Search](#) - [doi.wiley.com](#) - [ncbi.nlm.nih.gov](#) - [ncbi.nlm.nih.gov](#)

The search for Lps: 1993-1998

B Beutler, A Poltorak - Journal of Endotoxin Research, 2000 - [ingentaconnect.com](#)
 ... between the loci; D4Mit, designation of **unique** **sequence** markers isolated ... through-put **sequence** analysis of the region ... had become the distal border of the **contig**. ...
 Cited by 2 - [Web Search](#) - [ingentaconnect.com](#) - [ncbi.nlm.nih.gov](#) - [csa.com](#)

[book] Exploring Parasite Genomes

D Rollinson - 1999 - [print.google.com](#)
 ... A genome **sequence** lists the **unique** order of all nucleotides and their arrangement into coding and non-coding regions and is essentially a starting point for ...
 Cited by 1 - [Web Search](#) - [journals.cambridge.org](#) - [Library Search](#)

Fifth International Workshop on Human Chromosome 1 Mapping 1999

S Gregory, T Matise, B Schutte, P White - Cytogenet Cell Genet, 1999 - [content.karger.com](#)
 ... ing frames, characterized genes, **repetitive** elements, and ... This **unique** bundling procedure simultaneously refines gene ... d: Total **sequence** **contig** cov- erage for ...
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Characterizing the hemodynamic response: effects of presentation rate, sampling procedure, and the ...

FM Miezin, L Maccotta, JM Ollinger, SE Petersen, ... - Neuroimage, 2000 - [people.hnl.bcm.tmc.edu](#)
 ... A pillow and thermoplastic face **mask** were used to ... functional run, 128 sets of 16 **contig**- uous, 8-mm ... These equations are **repetitive** for uniformly (fixed) spaced ...
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Impact of memory systems on computer architecture and system organization.

RE Matick - IBM Systems Journal, 1986 - [research.ibm.com](#)
 Page 1. Impact of memory systems on computer architecture and system organization The largest part of computer architecture, in both ...
[View as HTML](#) - [Web Search](#) - [domino.research.ibm.com](#) - [csa.com](#)

Free Software in Molecular Biology for Macintosh and MS Windows Computers

D Gilbert - Bioinformatics Methods and Protocols (S. Misener and SA ..., 1999 - [iubio.bio.indiana.edu](#)
 ... a MS DOS program to identify common **repetitive** elements in ... also able to identify and **mask** vector **sequence** ... ClustalW multiple alignment, CAP **contig** assembly, tacg ...
 Cited by 2 - [Cached](#) - [Web Search](#) - [iubio.bio.indiana.edu](#) - [ncbi.nlm.nih.gov](#)

Expression of the human CFTR gene from episomal oriP-EBNA 1-YACs in mouse cells

D Huertas, S Howe, A McGuigan, C Huxley - Human Molecular Genetics, 2000 - [hmg.oupjournals.org](#)
 ... centromere (CEN), a yeast autonomously replicating **sequence** (ARS), a ... The **repetitive** GlyAla region (IR3 region) of EBNA-1 ... shown to be able to **mask** proteins from ...
 Cited by 12 - [Web Search](#) - [hmg.oupjournals.org](#) - [ingentaconnect.com](#) - [ncbi.nlm.nih.gov](#) - [all 5 versions »](#)

[book] Human Cytogenetic Cancer Markers

SR Wolman, S Sell - 1997 - [print.google.com](#)

... For many forms of human cancer, however, **unique** and tumor-specific chromosomal patterns have not yet been discerned, largely because of the extent and ...

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Oral Sessions

HJ Tanke, J Wiegant, RPM van Gijlswijk, V ... - Cytogenet Cell Genet, 1999 - [content.karger.com](#)

... With this approach a **telomere mask** is computed ... chromosome rearrangements are essentially **unique** to a ... EBI's nucleotide and protein **sequence** databases Claire ...

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[book] Colorectal Cancer Methods and Protocols: Methods and Protocols

SM Powell - 2000 - [print.google.com](#)

Page 1. Colorectal Cancer Page 2. METHODS IN MOLECULAR MEDICINE TM John M.

Walker, SERIES EDITOR 62. Parkinson's Disease: Methods and ...

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L2	22	mask and repetitive and contig and sequence and unique and @ad<"20020101"	USPAT	OR	OFF	2005/12/01 14:31
L3	9	mask and repetitive and contig and sequence and unique and @ad<"20000101"	USPAT	OR	OFF	2005/12/01 14:32
L4	7	mask and repetitive and contig and sequence and unique and @ad<"20000101" and database	USPAT	OR	OFF	2005/12/01 14:32